

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Yu Shuai

October 1, 2025

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Yu Shuai

October 1, 2025

- 1 Method
- 2 Results

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

└ Contents

Contents

- Method
- Results

- 1 Method
 - Symmetry-reduced non-intrusive trajectory-based ROM (SR-NiTROM)
- 2 Results

Contents

- Method
 - Symmetry-reduced non-intrusive trajectory-based ROM (SR-NiTROM)
- Results

Formulation of the SR-NiTROM

- Consider a shift-equivariant FOM with traveling solutions:

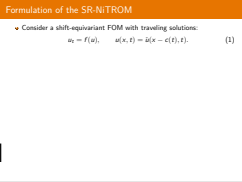
$$u_t = f(u), \quad u(x, t) = \hat{u}(x - c(t), t). \tag{1}$$

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

└ Method

└ Symmetry-reduced non-intrusive trajectory-based ROM (SR-NiTROM)



Formulation of the SR-NiTROM

- Consider a shift-equivariant FOM with traveling solutions:

$$u_t = f(u), \quad u(x, t) = \hat{u}(x - c(t), t). \tag{1}$$

- Suppose we have a collection of training snapshots $\{u(t_m)\}_{m=0}^{N_t-1}$, $u(t) \in \mathbb{R}^N$.

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

└ Method

└ Symmetry-reduced non-intrusive trajectory-based ROM (SR-NiTROM)

Formulation of the SR-NiTROM

Consider a shift-equivariant FOM with traveling solutions:
 $u_t = f(u), \quad u(x, t) = \hat{u}(x - c(t), t).$ (1)

Suppose we have a collection of training snapshots $\{u(t_m)\}_{m=0}^{N_t-1}$, $u(t) \in \mathbb{R}^N$.

Formulation of the SR-NiTROM

- Consider a shift-equivariant FOM with traveling solutions:

$$u_t = f(u), \quad u(x, t) = \hat{u}(x - c(t), t). \quad (1)$$

- Suppose we have a collection of training snapshots $\{u(t_m)\}_{m=0}^{N_t-1}$, $u(t) \in \mathbb{R}^N$.
- We seek to find $\Phi, \Psi \in \mathbb{R}^{n \times r}$, such that $\Phi(\Psi^\top \Phi)^{-1} \Psi^\top$ is a projection.

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Method

Symmetry-reduced non-intrusive trajectory-based ROM (SR-NiTROM)

Formulation of the SR-NiTROM

- Consider a shift-equivariant FOM with traveling solutions:
 $u_t = f(u), \quad u(x, t) = \hat{u}(x - c(t), t). \quad (1)$
- Suppose we have a collection of training snapshots $\{u(t_m)\}_{m=0}^{N_t-1}$, $u(t) \in \mathbb{R}^N$.
- We seek to find $\Phi, \Psi \in \mathbb{R}^{n \times r}$, such that $\Phi(\Psi^\top \Phi)^{-1} \Psi^\top$ is a projection.

Formulation of the SR-NiTROM

- Consider a shift-equivariant FOM with traveling solutions:

$$u_t = f(u), \quad u(x, t) = \hat{u}(x - c(t), t). \quad (1)$$

- Suppose we have a collection of training snapshots $\{u(t_m)\}_{m=0}^{N_t-1}$, $u(t) \in \mathbb{R}^N$.
- We seek to find $\Phi, \Psi \in \mathbb{R}^{n \times r}$, such that $\Phi(\Psi^\top \Phi)^{-1} \Psi^\top$ is a projection.
- From this projection operator, we can encode the FOM state with a low-dim representation $a(t) \in \mathbb{R}^r$:

$$\begin{aligned} a &= \Psi^\top u \\ \hat{u}_r &= \Phi(\Psi^\top \Phi)^{-1} a. \end{aligned} \quad (2)$$

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Method

Symmetry-reduced non-intrusive trajectory-based ROM (SR-NiTROM)

2025-10-01

Formulation of the SR-NiTROM

- Consider a shift-equivariant FOM with traveling solutions:
 $u_t = f(u), \quad u(x, t) = \hat{u}(x - c(t), t). \quad (1)$
- Suppose we have a collection of training snapshots $\{u(t_m)\}_{m=0}^{N_t-1}$, $u(t) \in \mathbb{R}^N$.
- We seek to find $\Phi, \Psi \in \mathbb{R}^{n \times r}$, such that $\Phi(\Psi^\top \Phi)^{-1} \Psi^\top$ is a projection.
- From this projection operator, we can encode the FOM state with a low-dim representation $a(t) \in \mathbb{R}^r$:
 $a = \Psi^\top u$
 $\hat{u}_r = \Phi(\Psi^\top \Phi)^{-1} a. \quad (2)$

Formulation of the SR-NiTROM

- Consider a shift-equivariant FOM with traveling solutions:

$$u_t = f(u), \quad u(x, t) = \hat{u}(x - c(t), t). \quad (1)$$

- Suppose we have a collection of training snapshots $\{u(t_m)\}_{m=0}^{N_t-1}$, $u(t) \in \mathbb{R}^N$.
- We seek to find $\Phi, \Psi \in \mathbb{R}^{n \times r}$, such that $\Phi(\Psi^\top \Phi)^{-1} \Psi^\top$ is a projection.
- From this projection operator, we can encode the FOM state with a low-dim representation $a(t) \in \mathbb{R}^r$:

$$a = \Psi^\top u$$

$$\hat{u}_r = \Phi(\Psi^\top \Phi)^{-1} a. \quad (2)$$

- The ROM dynamics is given in a symmetry-reduced form:

$$\dot{a}_i = A_{ij} a_j + B_{ijk} a_j a_k + \dot{c} M_{ij} a_j \quad (3a)$$

$$\dot{c} = - \frac{p_i a_i + Q_{ij} a_i a_j}{s_i a_i} \quad (3b)$$

$$M = \Psi^\top \partial_x \Phi (\Psi^\top \Phi)^{-1}, \quad s = \langle \partial_x \Phi (\Psi^\top \Phi)^{-1}, \partial_x u_0 \rangle \quad (3c)$$

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Method

Symmetry-reduced non-intrusive trajectory-based ROM (SR-NiTROM)

Formulation of the SR-NiTROM

- Consider a shift-equivariant FOM with traveling solutions:

$$u_t = f(u), \quad u(x, t) = \hat{u}(x - c(t), t). \quad (1)$$
- Suppose we have a collection of training snapshots $\{u(t_m)\}_{m=0}^{N_t-1}$, $u(t) \in \mathbb{R}^N$.
- We seek to find $\Phi, \Psi \in \mathbb{R}^{n \times r}$, such that $\Phi(\Psi^\top \Phi)^{-1} \Psi^\top$ is a projection.
- From this projection operator, we can encode the FOM state with a low-dim representation $a(t) \in \mathbb{R}^r$:

$$a = \Psi^\top u$$

$$\hat{u}_r = \Phi(\Psi^\top \Phi)^{-1} a. \quad (2)$$
- The ROM dynamics is given in a symmetry-reduced form:

$$\dot{a}_i = A_{ij} a_j + B_{ijk} a_j a_k + \dot{c} M_{ij} a_j \quad (3a)$$

$$\dot{c} = - \frac{p_i a_i + Q_{ij} a_i a_j}{s_i a_i} \quad (3b)$$

$$M = \Psi^\top \partial_x \Phi (\Psi^\top \Phi)^{-1}, \quad s = \langle \partial_x \Phi (\Psi^\top \Phi)^{-1}, \partial_x u_0 \rangle \quad (3c)$$

The optimization problem of SR-NiTROM

- The **trajectory-based** objective function:

$$J = \sum_{m=0}^{N_t-1} \|\hat{u}_r(t_m) - \hat{u}(t_m)\|^2 + \beta(c_r(t_m) - c(t_m))^2. \quad (4)$$

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

└ Method

└ Symmetry-reduced non-intrusive trajectory-based ROM (SR-NiTROM)

The optimization problem of SR-NiTROM

• The **trajectory-based** objective function:

$$J = \sum_{m=0}^{N_t-1} \|\hat{u}_r(t_m) - \hat{u}(t_m)\|^2 + \beta(c_r(t_m) - c(t_m))^2. \quad (4)$$

The optimization problem of SR-NiTROM

- The **trajectory-based** objective function:

$$J = \sum_{m=0}^{N_t-1} \|\hat{u}_r(t_m) - \hat{u}(t_m)\|^2 + \beta(c_r(t_m) - c(t_m))^2. \quad (4)$$

- $\beta = \gamma \sum_m \|\hat{u}(t_m)\|^2 / \sum_m (c(t_0) - c(t_m))^2$: relative weights.

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

└ Method

└ Symmetry-reduced non-intrusive trajectory-based ROM (SR-NiTROM)

The optimization problem of SR-NiTROM

• The **trajectory-based** objective function:

$$J = \sum_{m=0}^{N_t-1} \|\hat{u}_r(t_m) - \hat{u}(t_m)\|^2 + \beta(c_r(t_m) - c(t_m))^2. \quad (4)$$

• $\beta = \gamma \sum_m \|\hat{u}(t_m)\|^2 / \sum_m (c(t_0) - c(t_m))^2$: relative weights.

The optimization problem of SR-NiTROM

- The **trajectory-based** objective function:

$$J = \sum_{m=0}^{N_t-1} \|\hat{u}_r(t_m) - \hat{u}(t_m)\|^2 + \beta(c_r(t_m) - c(t_m))^2. \quad (4)$$

- $\beta = \gamma \sum_m \|\hat{u}(t_m)\|^2 / \sum_m (c(t_0) - c(t_m))^2$: relative weights.
- γ : hyperparameter.

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

└ Method

└ Symmetry-reduced non-intrusive trajectory-based ROM (SR-NiTROM)

The optimization problem of SR-NiTROM

• The **trajectory-based** objective function:

$$J = \sum_{m=0}^{N_t-1} \|\hat{u}_r(t_m) - \hat{u}(t_m)\|^2 + \beta(c_r(t_m) - c(t_m))^2. \quad (4)$$

• $\beta = \gamma \sum_m \|\hat{u}(t_m)\|^2 / \sum_m (c(t_0) - c(t_m))^2$: relative weights.

• γ : hyperparameter.

The optimization problem of SR-NiTROM

- The **trajectory-based** objective function:

$$J = \sum_{m=0}^{N_t-1} \|\hat{u}_r(t_m) - \hat{u}(t_m)\|^2 + \beta(c_r(t_m) - c(t_m))^2. \quad (4)$$

- $\beta = \gamma \sum_m \|\hat{u}(t_m)\|^2 / \sum_m (c(t_0) - c(t_m))^2$: relative weights.
- γ : hyperparameter.
- The unconstrained Lagrangian with multipliers:

$$L = \sum_{m=0}^{N_t-1} \left(\|\hat{u}_r(t_m) - \hat{u}(t_m)\|^2 + \beta(c_r(t_m) - c(t_m))^2 \right) + \int_{t_0}^{t_m} \lambda_m^\top (\dot{a} - Aa - B(a, a) - \dot{c}Ma) dt \quad (5)$$

$$+ \int_{t_0}^{t_m} \mu_m \left(\dot{c} + \frac{p_i a_i + Q_{ij} a_i a_j}{s_i a_i} \right) dt \quad (6)$$

$$+ \lambda_m(t_0)(a(t_0) - \Psi^\top \hat{u}(t_0))), \quad \lambda_m \in \mathbb{R}^r, \mu_m \in \mathbb{R}. \quad (7)$$

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Method

Symmetry-reduced non-intrusive trajectory-based ROM (SR-NiTROM)

The optimization problem of SR-NiTROM

- The **trajectory-based** objective function:

$$J = \sum_{m=0}^{N_t-1} \|\hat{u}_r(t_m) - \hat{u}(t_m)\|^2 + \beta(c_r(t_m) - c(t_m))^2. \quad (4)$$
 - $\beta = \gamma \sum_m \|\hat{u}(t_m)\|^2 / \sum_m (c(t_0) - c(t_m))^2$: relative weights.
 - γ : hyperparameter.
- The unconstrained Lagrangian with multipliers:

$$\begin{aligned} L = & \sum_{m=0}^{N_t-1} \left(\|\hat{u}_r(t_m) - \hat{u}(t_m)\|^2 + \beta(c_r(t_m) - c(t_m))^2 \right) \\ & + \int_{t_0}^{t_m} \lambda_m^\top (\dot{a} - Aa - B(a, a) - \dot{c}Ma) dt \\ & + \int_{t_0}^{t_m} \mu_m \left(\dot{c} + \frac{p_i a_i + Q_{ij} a_i a_j}{s_i a_i} \right) dt \\ & + \lambda_m(t_0)(a(t_0) - \Psi^\top \hat{u}(t_0)), \quad \lambda_m \in \mathbb{R}^r, \mu_m \in \mathbb{R}. \end{aligned} \quad (7)$$

- 1 Method
- 2 Results
 - Single trajectory: SR-NiTROM vs SR-Galerkin

Contents

- Method
- Results
 - Single trajectory: SR-NiTROM vs SR-Galerkin

- FOM: Kuramoto-Sivashinsky equation

$$u_t = -uu_x - u_{xx} - \nu u_{xxxx}, \quad x \in [0, 2\pi]. \tag{8}$$

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

- └ Results
 - └ Single trajectory: SR-NiTROM vs SR-Galerkin

● FOM: Kuramoto-Sivashinsky equation

$$u_t = -uu_x - u_{xx} - \nu u_{xxxx}, \quad x \in [0, 2\pi]. \tag{8}$$

Numerical details

- FOM: Kuramoto-Sivashinsky equation

$$u_t = -uu_x - u_{xx} - \nu u_{xxxxx}, \quad x \in [0, 2\pi]. \quad (8)$$

- $\nu = 4/87$ for traveling-wave patterns.

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Results

Single trajectory: SR-NiTROM vs SR-Galerkin

$$u_t = -u u_x - u_{xx} - v w_{xxxx}, \quad x \in [0, 2\pi]. \quad (8)$$

Numerical details

- FOM: Kuramoto-Sivashinsky equation

$$u_t = -uu_x - u_{xx} - \nu u_{xxxxx}, \quad x \in [0, 2\pi]. \quad (8)$$

- $\nu = 4/87$ for traveling-wave patterns.
- Periodic BCs, $N = 40$ Fourier modes, $\Delta t = 10^{-3}$.

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Results

└ Single trajectory: SR-NiTROM vs SR-Galerkin

$$u_2 = -u u_x - u_{xx} - v u_{xxx}, \quad x \in [0, 2\pi]. \quad (8)$$

- FOM: Kuramoto-Sivashinsky equation

$$u_t = -uu_x - u_{xx} - \nu u_{xxxx}, \quad x \in [0, 2\pi]. \quad (8)$$

- $\nu = 4/87$ for traveling-wave patterns.
- Periodic BCs, $N = 40$ Fourier modes, $\Delta t = 10^{-3}$.
- Sample interval: 10 timesteps between 2 adjacent snapshots.

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Results

Single trajectory: SR-NiTROM vs SR-Galerkin

Numerical details

• FOM: Kuramoto-Sivashinsky equation

$$u_t = -uu_x - u_{xx} - \nu u_{xxxx}, \quad x \in [0, 2\pi]. \quad (8)$$

- $\nu = 4/87$ for traveling-wave patterns.
- Periodic BCs, $N = 40$ Fourier modes, $\Delta t = 10^{-3}$.
- Sample interval: 10 timesteps between 2 adjacent snapshots.

- FOM: Kuramoto-Sivashinsky equation

$$u_t = -uu_x - u_{xx} - \nu u_{xxxx}, \quad x \in [0, 2\pi]. \quad (8)$$

- $\nu = 4/87$ for traveling-wave patterns.
 - Periodic BCs, $N = 40$ Fourier modes, $\Delta t = 10^{-3}$.
 - Sample interval: 10 timesteps between 2 adjacent snapshots.
- Optimization of the SR-NiTROM:

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Results

Single trajectory: SR-NiTROM vs SR-Galerkin

Numerical details

- FOM: Kuramoto-Sivashinsky equation

$$u_t = -uu_x - u_{xx} - \nu u_{xxxx}, \quad x \in [0, 2\pi]. \quad (8)$$

- $\nu = 4/87$ for traveling-wave patterns.
 - Periodic BCs, $N = 40$ Fourier modes, $\Delta t = 10^{-3}$.
 - Sample interval: 10 timesteps between 2 adjacent snapshots.
- Optimization of the SR-NiTROM:

- FOM: Kuramoto-Sivashinsky equation

$$u_t = -uu_x - u_{xx} - \nu u_{xxxx}, \quad x \in [0, 2\pi]. \quad (8)$$

- $\nu = 4/87$ for traveling-wave patterns.
 - Periodic BCs, $N = 40$ Fourier modes, $\Delta t = 10^{-3}$.
 - Sample interval: 10 timesteps between 2 adjacent snapshots.
- Optimization of the SR-NiTROM:
 - coordinate-descent method conjugate gradient optimizer for each subproblems.

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Results

Single trajectory: SR-NiTROM vs SR-Galerkin

Numerical details

- FOM: Kuramoto-Sivashinsky equation

$$u_t = -uu_x - u_{xx} - \nu u_{xxxx}, \quad x \in [0, 2\pi]. \quad (8)$$

- $\nu = 4/87$ for traveling-wave patterns.
 - Periodic BCs, $N = 40$ Fourier modes, $\Delta t = 10^{-3}$.
 - Sample interval: 10 timesteps between 2 adjacent snapshots.
- Optimization of the SR-NiTROM:
 - coordinate-descent method conjugate gradient optimizer for each subproblems.

- FOM: Kuramoto-Sivashinsky equation

$$u_t = -uu_x - u_{xx} - \nu u_{xxxx}, \quad x \in [0, 2\pi]. \quad (8)$$

- $\nu = 4/87$ for traveling-wave patterns.
 - Periodic BCs, $N = 40$ Fourier modes, $\Delta t = 10^{-3}$.
 - Sample interval: 10 timesteps between 2 adjacent snapshots.
- Optimization of the SR-NiTROM:
 - coordinate-descent method conjugate gradient optimizer for each subproblems.
 - 20 outer loops, 5 CG updates per outer loops.

$$u_t = -uu_x - u_{xx} - \nu u_{xxxx}, \quad x \in [0, 2\pi]. \quad (8)$$

- $\nu = 4/87$ for traveling-wave patterns.
 - Periodic BCs, $N = 40$ Fourier modes, $\Delta t = 10^{-3}$.
 - Sample interval: 10 timesteps between 2 adjacent snapshots.
- Optimization of the SR-NiTROM:
 - coordinate-descent method conjugate gradient optimizer for each subproblems.
 - 20 outer loops, 5 CG updates per outer loops.

- FOM: Kuramoto-Sivashinsky equation

$$u_t = -uu_x - u_{xx} - \nu u_{xxxx}, \quad x \in [0, 2\pi]. \quad (8)$$

- $\nu = 4/87$ for traveling-wave patterns.
- Periodic BCs, $N = 40$ Fourier modes, $\Delta t = 10^{-3}$.
- Sample interval: 10 timesteps between 2 adjacent snapshots.
- Optimization of the SR-NiTROM:
 - coordinate-descent method conjugate gradient optimizer for each subproblems.
 - 20 outer loops, 5 CG updates per outer loops.
 - Initial conditions: POD bases (capturing >99.5% energy) + Galerkin-projected tensors. (imitating the training result of the re-projected SR-OpInf ROM)

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Results

Single trajectory: SR-NiTROM vs SR-Galerkin

- FOM: Kuramoto-Sivashinsky equation

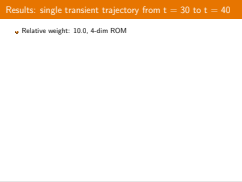
$$u_t = -uu_x - u_{xx} - \nu u_{xxxx}, \quad x \in [0, 2\pi]. \quad (8)$$

- $\nu = 4/87$ for traveling-wave patterns.
- Periodic BCs, $N = 40$ Fourier modes, $\Delta t = 10^{-3}$.
- Sample interval: 10 timesteps between 2 adjacent snapshots.
- Optimization of the SR-NiTROM:
 - coordinate-descent method conjugate gradient optimizer for each subproblems.
 - 20 outer loops, 5 CG updates per outer loops.
 - Initial conditions: POD bases (capturing >99.5% energy) + Galerkin-projected tensors. (imitating the training result of the re-projected SR-OpInf ROM)

Results: single transient trajectory from $t = 30$ to $t = 40$

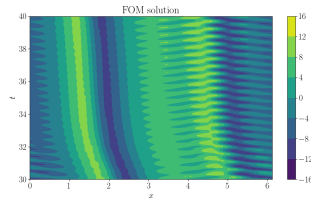
- Relative weight: 10.0, 4-dim ROM

2025-10-01
Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics
└ Results
└ Single trajectory: SR-NiTROM vs SR-Galerkin

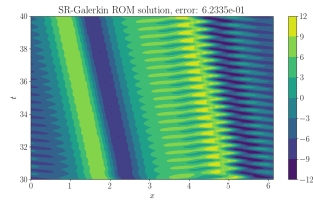


Results: single transient trajectory from $t = 30$ to $t = 40$

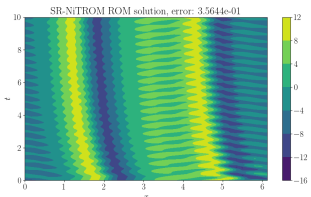
- Relative weight: 10.0, 4-dim ROM
- FOM vs SR-Galerkin vs SR-NiTROM



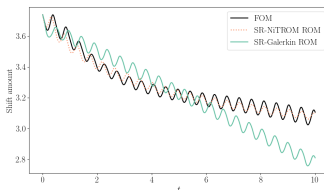
(a) FOM



(b) SR-Galerkin



(c) SR-NiTROM



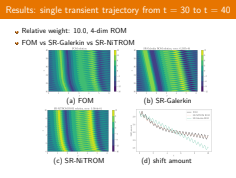
(d) shift amount

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Results

Single trajectory: SR-NiTROM vs SR-Galerkin



Results: multiple trajectories from perturbed initial conditions

- 9 trajectories, 7-dim ROM

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

- └ Results
 - └ Single trajectory: SR-NiTROM vs SR-Galerkin

Results: multiple trajectories from perturbed initial conditions

▼ 9 trajectories, 7-dim ROM

Results: multiple trajectories from perturbed initial conditions

- 9 trajectories, 7-dim ROM
- Initial conditions: post-transient solution snapshot + perturbations $u(t = 80) + \{0, \sin(x), \dots, \sin(4x), \cos(x), \dots, \cos(4x)\}$.

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

– Results

└ Single trajectory: SR-NiTROM vs SR-Galerkin

- 9 trajectories, 7-dim ROM
- Initial conditions: post-transient solution snapshot + perturbations $w(t=80) + \{0, \sin(x), \dots, \sin(4x), \cos(x), \dots, \cos(4x)\}$.

Results: multiple trajectories from perturbed initial conditions

- 9 trajectories, 7-dim ROM
- Initial conditions: post-transient solution snapshot + perturbations $u(t = 80) + \{0, \sin(x), \dots, \sin(4x), \cos(x), \dots, \cos(4x)\}$.
- Strategy A: 20 outer training loops (10 on bases, 10 on tensors), 5 CG updates per outer loop

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

└ Results

└ Single trajectory: SR-NiTROM vs SR-Galerkin

Results: multiple trajectories from perturbed initial conditions

- 9 trajectories, 7-dim ROM
- Initial conditions: post-transient solution snapshot + perturbations $u(t = 80) + \{0, \sin(x), \dots, \sin(4x), \cos(x), \dots, \cos(4x)\}$.
- Strategy A: 20 outer training loops (10 on bases, 10 on tensors), 5 CG updates per outer loop

Results: multiple trajectories from perturbed initial conditions

- 9 trajectories, 7-dim ROM
- Initial conditions: post-transient solution snapshot + perturbations $u(t = 80) + \{0, \sin(x), \dots, \sin(4x), \cos(x), \dots, \cos(4x)\}$.
- Strategy A: 20 outer training loops (10 on bases, 10 on tensors), 5 CG updates per outer loop
- Strategy B: 10 outer training loops on tensors only with fixed POD bases

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Results

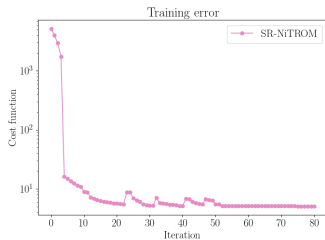
Single trajectory: SR-NiTROM vs SR-Galerkin

Results: multiple trajectories from perturbed initial conditions

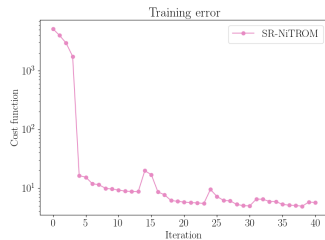
- 9 trajectories, 7-dim ROM
- Initial conditions: post-transient solution snapshot + perturbations $u(t = 80) + \{0, \sin(x), \dots, \sin(4x), \cos(x), \dots, \cos(4x)\}$.
- Strategy A: 20 outer training loops (10 on bases, 10 on tensors), 5 CG updates per outer loop
- Strategy B: 10 outer training loops on tensors only with fixed POD bases

Results: multiple trajectories from perturbed initial conditions

- 9 trajectories, 7-dim ROM
- Initial conditions: post-transient solution snapshot + perturbations $u(t=80) + \{0, \sin(x), \dots, \sin(4x), \cos(x), \dots, \cos(4x)\}$.
- Strategy A: 20 outer training loops (10 on bases, 10 on tensors), 5 CG updates per outer loop
- Strategy B: 10 outer training loops on tensors only with fixed POD bases
- Training loss: **not too much difference on the training set.**



(a) Strategy A



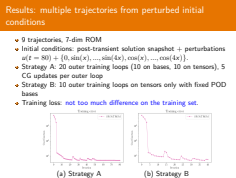
(b) Strategy B

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

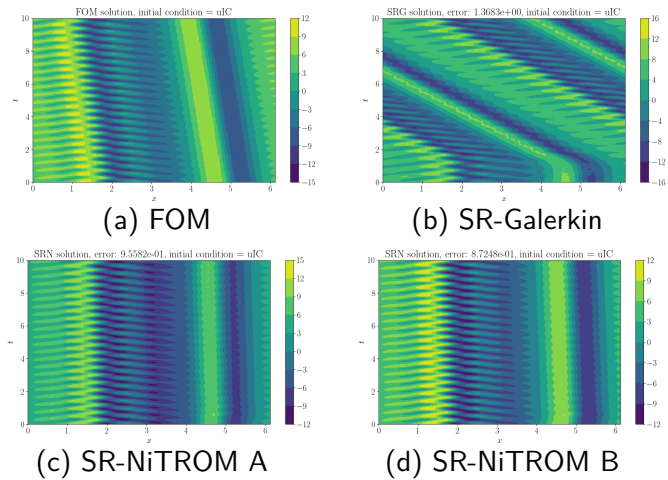
Results

Single trajectory: SR-NiTROM vs SR-Galerkin



Results: multiple trajectories from perturbed initial conditions

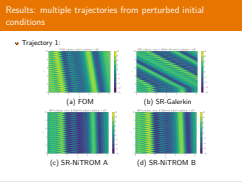
- Trajectory 1:



2025-10-01

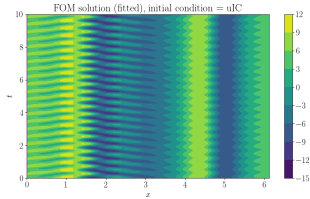
Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

- Results
 - Single trajectory: SR-NiTROM vs SR-Galerkin

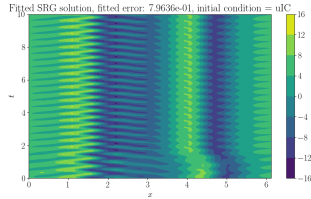


Results: multiple trajectories from perturbed initial conditions

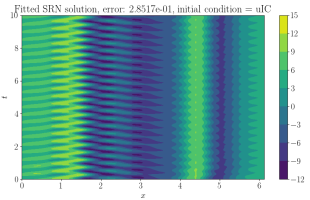
- Trajectory 1:



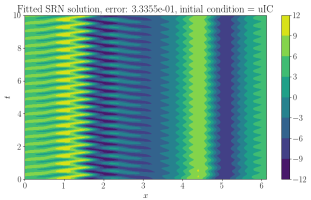
(a) FOM



(b) SR-Galerkin



(c) SR-NiTROM A

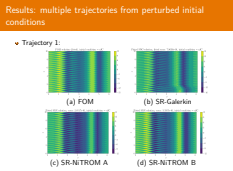


(d) SR-NiTROM B

2025-10-01

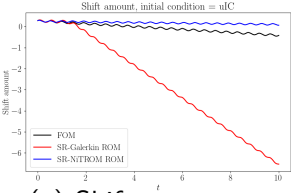
Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

- Results
 - Single trajectory: SR-NiTROM vs SR-Galerkin

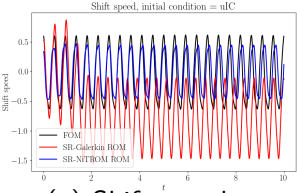


Results: multiple trajectories from perturbed initial conditions

- Trajectory 1, shift amount and shift speed:



(a) Shift amounts

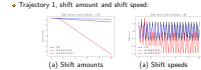


(a) Shift speeds

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

- Results
 - Single trajectory: SR-NiTROM vs SR-Galerkin



Conclusions:

- For the reconstruction of a single training trajectory **including transient**, SR-NiTROM outperforms SR-Galerkin ROM (and SR-OpInf of course).

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Results

└ Single trajectory: SR-NiTROM vs SR-Galerkin

- For the reconstruction of a single training trajectory **including transient**, SR-NiTROM outperforms SR-Galerkin ROM (and SR-OpInf of course).

Conclusions:

- For the reconstruction of a single training trajectory [including transient](#), SR-NiTROM outperforms SR-Galerkin ROM (and SR-OpInf of course).
- For the reconstruction of multiple transient trajectories, we find that:

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

└ Results

└ Single trajectory: SR-NiTROM vs SR-Galerkin

Conclusions:

- For the reconstruction of a single training trajectory [including transient](#), SR-NiTROM outperforms SR-Galerkin ROM (and SR-OpInf of course).
- For the reconstruction of multiple transient trajectories, we find that:

Conclusions:

- For the reconstruction of a single training trajectory **including transient**, SR-NiTROM outperforms SR-Galerkin ROM (and SR-OpInf of course).
- For the reconstruction of multiple transient trajectories, we find that:
 - SR-NiTROM gives better approximation of template-aligned snapshots and shift amounts than the SR-Galerkin ROM.

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

- Results

Single trajectory: SR-NiTROM vs SR-Galerkin

- For the reconstruction of a single training trajectory including transient, SR-NiTROM outperforms SR-Galerkin ROM (and SR-Opnf of course).
- For the reconstruction of multiple transient trajectories, we find that:
 - SR-NiTROM gives better approximation of template-aligned snapshots and shift amounts than the SR-Galerkin ROM.

Conclusions:

- For the reconstruction of a single training trajectory **including transient**, SR-NiTROM outperforms SR-Galerkin ROM (and SR-OpInf of course).
- For the reconstruction of multiple transient trajectories, we find that:
 - SR-NiTROM gives better approximation of template-aligned snapshots and shift amounts than the SR-Galerkin ROM.
 - It's better to optimize both the bases and the tensors to minimize our loss function.

2025-10-01

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Results

-Single trajectory: SR-NiTROM vs SR-Galerkin

- For the reconstruction of a single training trajectory including transient, SR-NITROM outperforms SR-Galerkin ROM (and SR-OpIn of course).
- For the reconstruction of multiple transient trajectories, we find that:
 - SR-NITROM gives better approximation of template-aligned snapshots and shift amounts than the SR-Galerkin ROM.
 - It's better to optimize both the bases and the tensors to minimize our loss function.

Conclusions:

- For the reconstruction of a single training trajectory **including transient**, SR-NiTROM outperforms SR-Galerkin ROM (and SR-OpInf of course).
- For the reconstruction of multiple transient trajectories, we find that:
 - SR-NiTROM gives better approximation of template-aligned snapshots and shift amounts than the SR-Galerkin ROM.
 - It's better to optimize both the bases and the tensors to minimize our loss function.
 - **However, SR-NiTROM with only trained tensors and POD bases attains the least reconstruction error of the raw snapshots.**

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Results

Single trajectory: SR-NiTROM vs SR-Galerkin

Conclusions:

- For the reconstruction of a single training trajectory **including transient**, SR-NiTROM outperforms SR-Galerkin ROM (and SR-OpInf of course).
- For the reconstruction of multiple transient trajectories, we find that:
 - SR-NiTROM gives better approximation of template-aligned snapshots and shift amounts than the SR-Galerkin ROM.
 - It's better to optimize both the bases and the tensors to minimize our loss function.
 - **However, SR-NiTROM with only trained tensors and POD bases attains the least reconstruction error of the raw snapshots.**

Conclusions:

- For the reconstruction of a single training trajectory **including transient**, SR-NiTROM outperforms SR-Galerkin ROM (and SR-OpInf of course).
- For the reconstruction of multiple transient trajectories, we find that:
 - SR-NiTROM gives better approximation of template-aligned snapshots and shift amounts than the SR-Galerkin ROM.
 - It's better to optimize both the bases and the tensors to minimize our loss function.
 - **However, SR-NiTROM with only trained tensors and POD bases attains the least reconstruction error of the raw snapshots.**
 - Why: our loss function = error in aligned snapshots + error in shift amounts, not error in raw snapshots.

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Results

Single trajectory: SR-NiTROM vs SR-Galerkin

Conclusions:

- For the reconstruction of a single training trajectory **including transient**, SR-NiTROM outperforms SR-Galerkin ROM (and SR-OpInf of course).
- For the reconstruction of multiple transient trajectories, we find that:
 - SR-NiTROM gives better approximation of template-aligned snapshots and shift amounts than the SR-Galerkin ROM.
 - It's better to optimize both the bases and the tensors to minimize our loss function.
 - **However, SR-NiTROM with only trained tensors and POD bases attains the least reconstruction error of the raw snapshots.**
 - Why: our loss function = error in aligned snapshots + error in shift amounts, not error in raw snapshots.

Conclusions:

- For the reconstruction of a single training trajectory **including transient**, SR-NiTROM outperforms SR-Galerkin ROM (and SR-OpInf of course).
- For the reconstruction of multiple transient trajectories, we find that:
 - SR-NiTROM gives better approximation of template-aligned snapshots and shift amounts than the SR-Galerkin ROM.
 - It's better to optimize both the bases and the tensors to minimize our loss function.
 - **However, SR-NiTROM with only trained tensors and POD bases attains the least reconstruction error of the raw snapshots.**
 - Why: our loss function = error in aligned snapshots + error in shift amounts, not error in raw snapshots.
 - Trade-off: we may want to switch to error in raw snapshots, but then the optimizer doesn't know if the error comes from mismatch of aligned profiles or shift amounts.

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Results

Single trajectory: SR-NiTROM vs SR-Galerkin

Conclusions:

- For the reconstruction of a single training trajectory **including transient**, SR-NiTROM outperforms SR-Galerkin ROM (and SR-OpInf of course).
- For the reconstruction of multiple transient trajectories, we find that:
 - SR-NiTROM gives better approximation of template-aligned snapshots and shift amounts than the SR-Galerkin ROM.
 - It's better to optimize both the bases and the tensors to minimize our loss function.
 - **However, SR-NiTROM with only trained tensors and POD bases attains the least reconstruction error of the raw snapshots.**
 - Why: our loss function = error in aligned snapshots + error in shift amounts, not error in raw snapshots.
 - Trade-off: we may want to switch to error in raw snapshots, but then the optimizer doesn't know if the error comes from mismatch of aligned profiles or shift amounts.

Conclusions:

- For the reconstruction of a single training trajectory **including transient**, SR-NiTROM outperforms SR-Galerkin ROM (and SR-OpInf of course).
- For the reconstruction of multiple transient trajectories, we find that:
 - SR-NiTROM gives better approximation of template-aligned snapshots and shift amounts than the SR-Galerkin ROM.
 - It's better to optimize both the bases and the tensors to minimize our loss function.
 - **However, SR-NiTROM with only trained tensors and POD bases attains the least reconstruction error of the raw snapshots.**
 - Why: our loss function = error in aligned snapshots + error in shift amounts, not error in raw snapshots.
 - Trade-off: we may want to switch to error in raw snapshots, but then the optimizer doesn't know if the error comes from mismatch of aligned profiles or shift amounts.
 - This new loss function is reasonable since small shift mismatch can lead to large error in raw snapshots.

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Results

Single trajectory: SR-NiTROM vs SR-Galerkin

2025-10-01

Conclusions:

- For the reconstruction of a single training trajectory **including transient**, SR-NiTROM outperforms SR-Galerkin ROM (and SR-OpInf of course).
- For the reconstruction of multiple transient trajectories, we find that:
 - SR-NiTROM gives better approximation of template-aligned snapshots and shift amounts than the SR-Galerkin ROM.
 - It's better to optimize both the bases and the tensors to minimize our loss function.
 - **However, SR-NiTROM with only trained tensors and POD bases attains the least reconstruction error of the raw snapshots.**
 - Why: our loss function = error in aligned snapshots + error in shift amounts, not error in raw snapshots.
 - Trade-off: we may want to switch to error in raw snapshots, but then the optimizer doesn't know if the error comes from mismatch of aligned profiles or shift amounts.
 - This new loss function is reasonable since small shift mismatch can lead to large error in raw snapshots.

Conclusions:

- For the reconstruction of a single training trajectory **including transient**, SR-NiTROM outperforms SR-Galerkin ROM (and SR-OpInf of course).
- For the reconstruction of multiple transient trajectories, we find that:
 - SR-NiTROM gives better approximation of template-aligned snapshots and shift amounts than the SR-Galerkin ROM.
 - It's better to optimize both the bases and the tensors to minimize our loss function.
 - **However, SR-NiTROM with only trained tensors and POD bases attains the least reconstruction error of the raw snapshots.**
 - Why: our loss function = error in aligned snapshots + error in shift amounts, not error in raw snapshots.
 - Trade-off: we may want to switch to error in raw snapshots, but then the optimizer doesn't know if the error comes from mismatch of aligned profiles or shift amounts.
 - This new loss function is reasonable since small shift mismatch can lead to large error in raw snapshots.
- To-dos: test our SR-NiTROM on unseen trajectories. Compute the obliqueness of projection.

Non-intrusive model reduction of shift-equivariant systems via data-driven projection and reduced dynamics

Results

Single trajectory: SR-NiTROM vs SR-Galerkin

Conclusions:

- For the reconstruction of a single training trajectory **including transient**, SR-NiTROM outperforms SR-Galerkin ROM (and SR-OpInf of course).
- For the reconstruction of multiple transient trajectories, we find that:
 - SR-NiTROM gives better approximation of template-aligned snapshots and shift amounts than the SR-Galerkin ROM.
 - It's better to optimize both the bases and the tensors to minimize our loss function.
 - **However, SR-NiTROM with only trained tensors and POD bases attains the least reconstruction error of the raw snapshots.**
 - Why: our loss function = error in aligned snapshots + error in shift amounts, not error in raw snapshots.
 - Trade-off: we may want to switch to error in raw snapshots, but then the optimizer doesn't know if the error comes from mismatch of aligned profiles or shift amounts.
 - This new loss function is reasonable since small shift mismatch can lead to large error in raw snapshots.
- To-dos: test our SR-NiTROM on unseen trajectories. Compute the obliqueness of projection.